

1/9

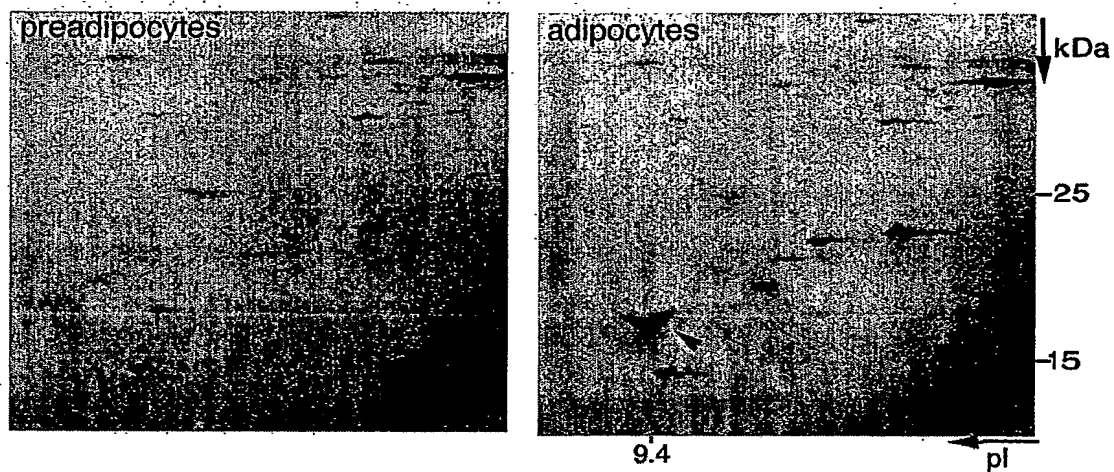
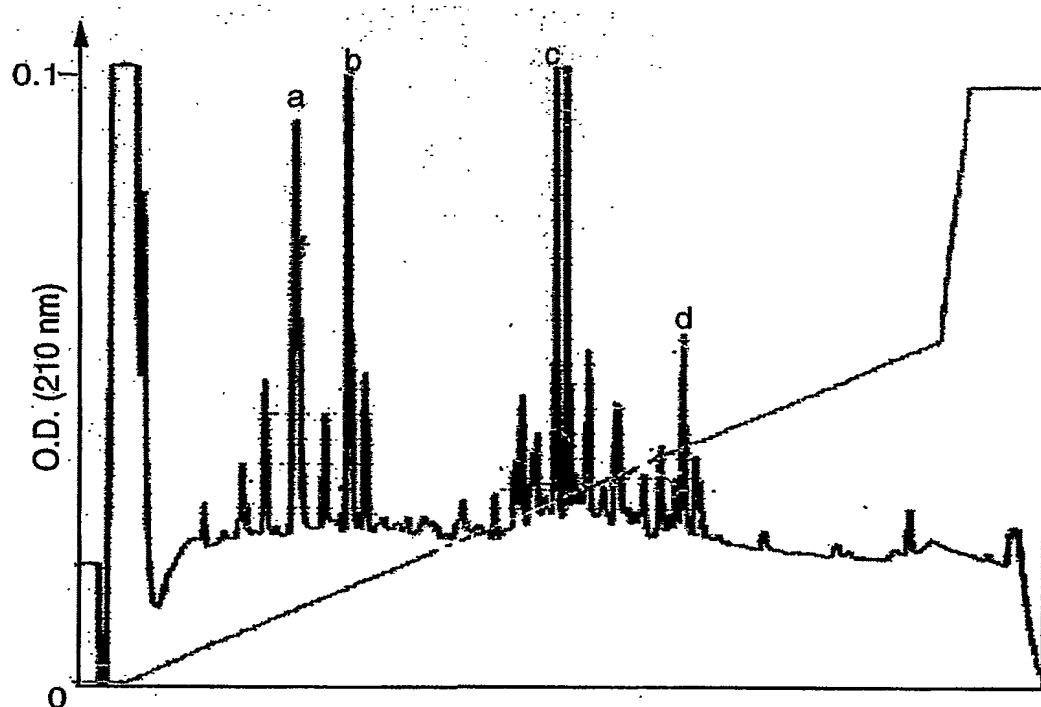


Figure 1

2/9



Farfctions	Amino acid sequences
a	LQQTNC PK
b	GTEPELSETQR
c	CLLISLALWLGTVGTR
d	HPPVQLAFQEIGVDR

Figure 2

3/9

SEQUENCE ID NO: 1

MKGILSTALWECTVGH RGTEPELSETQRRSLQVALEEFHKHPPVQLA  
 FQBIGVDRAEEVLFSAGTFVRLEFKLQQTNPCKDWKKPECTIKPNGR  
 RRKCLACIKMDPKGKILGRIVHCPILKQGPQDPQELQCIKIAQAGEDP  
 HGYFLPGQFAFSRALRTK

B.

SS (1-17)

Potential cystatin like  
domain (32-108)Domain with unknown  
function (109-162)

C.

Adipocyspin

LQVLEFHHPVQLAFQEGDR EEVLFSAGTFVR EFKLQQNSPKDWKKP-T  
 90

Cystatin C

ADFVAHYNKSNDLYELRVKVS KSQVV-AGNNYYKVEVGEKLSPTDLNP 78

Query: 91

IKPNRRRK LACIKMD 108

Sbjct: 79

FLEQEEGFDFEVWEK 96

Figure 3